

Figure 1

Sequence alignment of mouse Serca 1, 2 and 3 protein.

5	Serca1a	1	MEAAHSKSTEECLSYFGVSETTGLTPDQVKRHLKYGPNELPAEEGKSLWELVVEQFEDL
	Serca2a	1	..N..T.TV..V.GH..N.S...SLE..KLK.RW.S.....T.L...I.....
	Serca2b	1	..N..T.TV..V.GH..N.S...SLE..KLK.RW.S.....T.L...I.....
	Serca3a	1	..E..LL.AADV.RR.S.TAEG..SLE..TDAR.R.....T.....
	Serca3b	1	..E..LL.AADV.RR.S.TAEG..SLE..TDAR.R.....T.....
10	Serca3c	1	..E..LL.AADV.RR.S.TAEG..SLE..TDAR.R.....T.....
	Serca1a	61	LVRILLAAACISFVLAWFEEGEETVTAFVEPFVILLILIANAIVGWQERNAENAIEALK
	Serca2a	61	.....I.....V.....
	Serca2b	61	.....I.....V.....
15	Serca3a	61	.....LV.....T.....L..M..V.....S.....
	Serca3b	61	.....LV.....T.....L..M..V.....S.....
	Serca3c	61	.....LV.....T.....L..M..V.....S.....
	Serca1a	121	EYEPEMGKVYRADRKSVQRIKARDIVPGDIVEVAVGDKVPADIRILSIKSTTLRVDQSL
20	Serca2a	121	.....Q.....K.....I.....LT.....
	Serca2b	121	.....Q.....K.....I.....LT.....
	Serca3a	121	.....I.S...G....R.....L.LIE.....
	Serca3b	121	.....I.S...G....R.....L.LIE.....
	Serca3c	121	.....I.S...G....R.....L.LIE.....
25	Serca1a	181	TGESVSVIKHTDPVPDPRAVNQDKKNMLFSGTNIAAGKAVGIVATTGVSTEIGKIRDQMA
	Serca2a	181	.....M.V.VA...N.....E.V
	Serca2b	181	.....M.V.VA...N.....E.V
	Serca3a	181	.....T....AI.....S...L.VAVA..LQ..L....S...
30	Serca3b	181	.....T....AI.....S...L.VAVA..LQ..L....S...
	Serca3c	181	.....T....AI.....S...L.VAVA..LQ..L....S...
	Serca1a	241	ATEQDKTPLQQKLDEFGEQLSKVISLICVAVWLINIGHFNDPVHGGSWFRGAIYYFKIAV
	Serca2a	241	....ER.....I..I.....I.....
35	Serca2b	241	....ER.....I..I.....I.....
	Serca3a	241	.V.PER....R...HA..V.....V.....A..A....L..V.....
	Serca3b	241	.V.PER....R...HA..V.....V.....A..A....L..V.....
	Serca3c	241	.V.PER....R...HA..V.....V.....A..A....L..V.....
40	Serca1a	301	ALAVAAIPEGLPAVITTCIALGTRRMAKKNAIVRSLPSVETLGCTSVICSDKTGTLTTNQ
	Serca2a	301	.....
	Serca2b	301	.....
	Serca3a	301	.....R.....
	Serca3b	301	.....R.....
45	Serca3c	301	.....R.....
	Serca1a	361	MSVCKMFIIDKVDGDVCSLNEFSITGSTYAPEGEVLKNDKPVRAGQYDGLVELATICALC
	Serca2a	361	....R...L...E..T.....I...Q.D....KCH.....
	Serca2b	361	....R...L...E..T.....I...Q.D....KCH.....
50	Serca3a	361	....R..VVAEAEAGT.R.H..T.S.T..T....RQGEQ..C..F.....
	Serca3b	361	....R..VVAEAEAGT.R.H..T.S.T..T....RQGEQ..C..F.....
	Serca3c	361	....R..VVAEAEAGT.R.H..T.S.T..T....RQGEQ..C..F.....
	Serca1a	421	NDSSLDFNETKGVYEVGEATETALTTLVEKMNVFNTTEVRSLSKVERANACNSVIQLMK
55	Serca2a	421	...A..Y..A.....C.....D..LKG..I.....K....
	Serca2b	421	...A..Y..A.....C.....D..LKG..I.....K....
	Serca3a	421	...A..Y..A.....C.....D..DLKG..R...G.....K...R
	Serca3b	421	...A..Y..A.....C.....D..DLKG..R...G.....K...R
	Serca3c	421	...A..Y..A.....C.....D..DLKG..R...G.....K...R

Serca1a	481	KEFTLEFSRDRKSMCSVYCSPAKSSRAAVGNKMFVKGAPEGVIDRCNYVRVGTTRVPLTGP	
Serca2a	481	.....T.N.P..TSMS-.....THI...S.K..M.PG	
Serca2b	481	.....T.N.P..TSMS-.....THI...S.K..M.PG	
Serca3a	481	.....T.TRADPKVQ.S.....S..E..SS...SRTA..STT	
5	Serca3b	481	.....T.TRADPKVQ.S.....S..E..SS...SRTA..STT
	Serca3c	481	.....T.TRADPKVQ.S.....S..E..SS...SRTA..STT
Serca1a	541	VKEKIMSVIKEWGTGRDTLRLALATRDTPPKREEMVLDDSAKFMNEYEMDLTFVGVVGML	
Serca2a	540	..Q.....R...S.S.....H.N.L.....H.E...N.IK..TN.....C....	
10	Serca2b	540	..Q.....R...S.S.....H.N.L.....H.E...N.IK..TN.....C....
	Serca3a	541	SR.H.LAK.RD..S.S.....RK.D.H...CSR.VQ..T.....C....
	Serca3b	541	SR.H.LAK.RD..S.S.....RK.D.H...CSR.VQ..T.....C....
	Serca3c	541	SR.H.LAK.RD..S.S.....RK.D.H...CSR.VQ..T.....C....
15	Serca1a	601	DPPRKEVTGSIQLCRDAGIRVIMITGDNKGTAIAICRRIGIFSENEEVTDRAYTGREFDD
	Serca2a	600	....I..AS.VK..Q.....V.....GQD.D..SK.F.....E
	Serca2b	600	....I..AS.VK..Q.....V.....GQD.D..SK.F.....E
	Serca3a	601	....P..AAC.TR.SR....V.....V.....L..GDT.D.LGK.....
	Serca3b	601	....P..AAC.TR.SR....V.....V.....L..GDT.D.LGK.....
20	Serca3c	601	....P..AAC.TR.SR....V.....V.....L..GDT.D.LGK.....
Serca1a	661	LPLAEQREACRRACCFARVEPSHKSIVEYLQSYDEITAMTGDGVNDAPALKKAEIGIAM	
Serca2a	660	.SPSA..D..LN.R.....F...F.....S.....	
Serca2b	660	.SPSA..D..LN.R.....F...F.....S.....	
25	Serca3a	661	.SPEQ..Q...T.R.....A..R..N..FN.....
	Serca3b	661	.SPEQ..Q...T.R.....A..R..N..FN.....
	Serca3c	661	.SPEQ..Q...T.R.....A..R..N..FN.....
Serca1a	721	GSGTAVAKTASEMVLADDNFSTIVAAVEGRAIYNNMKQFIRYLISSNVGEVVCIFLTAA	
30	Serca2a	720	.....
	Serca2b	720	.....
	Serca3a	721	....S.A....S....AS.....I
	Serca3b	721	....S.A....S....AS.....I
	Serca3c	721	....S.A....S....AS.....I
35	Serca1a	781	LGLPEALIPVQLLWVNLVTDGLPATALGFNPPDLDIMDRPPRSPKEPLISGWLFFRYMAI
	Serca2a	780	..F.....NK..N.....L..
	Serca2b	780	..F.....NK..N.....L..
	Serca3a	781	.....EK..N.R.A.....L..
40	Serca3b	781	.....EK..N.R.A.....L..
	Serca3c	781	.....EK..N.R.A.....L..
Serca1a	841	GGYVGAATVGAAAWFLYAEDGPHVSYHQLTHFMQCTEHNPEFDGLCEVFEAPEPMTMA	
Serca2a	840	.C.....IA.DG..R..FY..S..L..K.D..D..V..AI..S.Y.....	
45	Serca2b	840	.C.....IA.DG..R..FY..S..L..K.D..D..V..AI..S.Y.....
	Serca3a	841	.V...L..A..T.....DAE..Q.TFY..RN.LK.S.D..L.A.I..K..SRF.T...
	Serca3b	841	.V...L..A..T.....DAE..Q.TFY..RN.LK.S.D..L.A.I..K..SRF.T...
	Serca3c	841	.V...L..A..T.....DAE..Q.TFY..RN.LK.S.D..L.A.I..K..SRF.T...
50	Serca1a	901	LSVLVTIEMCNALNSLSENQSLLRMPPWVNIWLLGSICLSMSLHFLILYVDPLPMIFKLR
	Serca2a	900	.....E.....V.....E...L..QIT
	Serca2b	900	.....E.....V.....E...L..QIT
	Serca3a	901	.....V.....L.P....AVVM..A.....L.P...L..QVT
	Serca3b	901	.....V.....L.P....AVVM..A.....L.P...L..QVT
55	Serca3c	901	.....V.....L.P....AVVM..A.....L.P...L..QVT

Sercala 961 ALDFTQWLMVLKISLPVIGLDELLKFIARNYLEG  
Serc2a 960 P.NL.....LM..T....V.....QPAILE  
Serc2b 960 P.NL.....LM..T....V.....QPGKECVQPATKSSCSLSACTDGISWP  
Serc3a 961 P.SGR..GV..QM....L...A..YLS..HMDEKKDLK  
5 Serc3b 961 P.SGR..GV..QM....L...A..YLS..HMD.VLGTFMQARSRQLPTTSRTPYHTGKK  
Serc3c 961 P.SGR..GV..QM....L...A..YLS..HMD.VLGTFMQARSRQLPTTSRTPYHTGLA

Serca2b 1020 FVLLIMPLVVWVYSTDNFSDMFWS  
10 Serc3b 1021 GPEVNPGSRGESPVWPSD  
Serc3c 1021 SWKKRT

Figure 2

Sequence similarity of Serca2 proteins in mammalian species

5	Mouse_2a	1	MENAHTKTVEEVLGHFGVNESTGLSLEQVKKLKERWGSNELPAEEGKTLL ELVIEQFEDL
	Mouse_2b	1	.....
	Rat_2b	1	.....
	Rat_2a	1	.....
	Dog_2a	1	.....
	Cat_2a	1	.....Y.....
10	Pig_2a	1	.....
	Pig_2b	1	.....
	Human_2b	1	.....
	Human_2c	1	.....
	Human_2a	1	.....
15	Rabbit_2a	1	.....
	Rabbit_2b	1	.....
	Mouse_2a	61	LVRILLLAACISFVLAWFEEGEETITAFVEPFVILLILVANAIIVGVWQERNA ENATEALK
	Mouse_2b	61	.....
20	Rat_2b	61	.....
	Rat_2a	61	.....
	Dog_2a	61	.....
	Cat_2a	61	.....
	Pig_2a	61	.....
25	Pig_2b	61	.....
	Human_2b	61	.....
	Human_2c	61	.....
	Human_2a	61	.....
	Rabbit_2a	61	.....
30	Rabbit_2b	61	.....
	Mouse_2a	121	EYEPEMGKVYRQDRKSVQRRIKAKDIVPGDIVEIAVGDKVPADIRLTSIK STTLLRVDQSI
	Mouse_2b	121	.....
	Rat_2b	121	.....
35	Rat_2a	121	.....
	Dog_2a	121	.....
	Cat_2a	121	.....
	Pig_2a	121	.....
	Pig_2b	121	.....
40	Human_2b	121	.....
	Human_2c	121	.....
	Human_2a	121	.....
	Rabbit_2a	121	.....
	Rabbit_2b	121	.....
45	Mouse_2a	181	TGESVSVIKHTDPVPDPRAVNQDKKNMLFSGTNIAAGKAMGVV VATGVNTEIGKIRDEM
	Mouse_2b	181	.....
	Rat_2b	181	.....
	Rat_2a	181	.....
50	Dog_2a	181	.....
	Cat_2a	181	.....
	Pig_2a	181	.....
	Pig_2b	181	.....
	Human_2b	181	.....
55	Human_2c	181	.....
	Human_2a	181	.....
	Rabbit_2a	181	.....
	Rabbit_2b	181	.....

	Mouse_2a	241	ATEQERTPLQQKLDEFGEQLSKVISLICIAVWIINIGHFNDPVHGGSWIRGAIYYFKIAV
	Mouse_2b	241	.....
5	Rat_2b	241	.....
	Rat_2a	241	.....
	Dog_2a	241	.....
	Cat_2a	241	.....
	Pig_2a	241	.....
	Pig_2b	241	.....
10	Human_2b	241	.....
	Human_2c	241	.....
	Human_2a	241	.....
	Rabbit_2a	241	.....
	Rabbit_2b	241	.....
15	Mouse_2a	301	ALAVAAIPEGLPAVITTCALGTRRMAKKNAIVRSLPSVETLGCTSVICSDKTGTLTTNQ
	Mouse_2b	301	.....
	Rat_2b	301	.....
	Rat_2a	301	.....
20	Dog_2a	301	.....
	Cat_2a	301	.....
	Pig_2a	301	.....
	Pig_2b	301	.....
	Human_2b	301	.....
25	Human_2c	301	.....
	Human_2a	301	.....
	Rabbit_2a	301	.....
	Rabbit_2b	301	.....
30	Mouse_2a	361	MSVCRMFIIDKVEGDTCSLNEFSITGSTYAPIGEVQKDDKPVKCHQYDGLVELATICALC
	Mouse_2b	361	.....
	Rat_2b	361	.....T.....
	Rat_2a	361	.....T.....
	Dog_2a	361	.....R.....S.....T.....H.....
35	Cat_2a	361	.....T.....H.....
	Pig_2a	361	.....T.....H.....
	Pig_2b	361	.....T.....H.....
	Human_2b	361	.....R.....T.....H.....N.....
	Human_2c	361	.....R.....T.....H.....N.....
40	Human_2a	361	.....R.....T.....H.....N.....
	Rabbit_2a	361	.....D.....T.....H.....
	Rabbit_2b	361	.....D.....T.....H.....
	Mouse_2a	421	NDSALDYNEAKGVYEVGEATETALTCLVEKMNVDTELKGLSKIERANACNSVIKQLMK
45	Mouse_2b	421	.....
	Rat_2b	421	.....
	Rat_2a	421	.....
	Dog_2a	421	.....
	Cat_2a	421	.....K.F.....
50	Pig_2a	421	.....
	Pig_2b	421	.....
	Human_2b	421	.....
	Human_2c	421	.....
	Human_2a	421	.....
55	Rabbit_2a	421	.....
	Rabbit_2b	421	.....

	Mouse_2a	481	KEFTLEFSRDRKSMHSVYCTPNKPSRTSMSKMFVKGAPEGVIDRCTHIRVGSTKVPMTPGV
	Mouse_2b	481	.....
	Rat_2b	481	.....
5	Rat_2a	481	.....
	Dog_2a	481	.....
	Cat_2a	481	.....
	Pig_2a	481	.....
	Pig_2b	481	.....
	Human_2b	481	.....
10	Human_2c	481	.....
	Human_2a	481	.....
	Rabbit_2a	481	.....
	Rabbit_2b	481	.....
15	Mouse_2a	541	KQKIMSVIREWGSGSDTLRCLALATHDNPLKREEMHLEDSANFIKYETNLT
	Mouse_2b	541	FVGCVGMLD
	Rat_2b	541	.....
	Rat_2a	541	.....
20	Dog_2a	541	.....V.....
	Cat_2a	541	.....V.....
	Pig_2a	541	.....
	Pig_2b	541	.....
	Human_2b	541	.....
25	Human_2c	541	.....
	Human_2a	541	.....
	Rabbit_2a	541	.....
	Rabbit_2b	541	.....
30	Mouse_2a	601	PPRIEVASSVKLCRQAGIRVIMITGDNKGTAVAICRRIGIFGQDEDVTSKAFTGREFDEL
	Mouse_2b	601	.....
	Rat_2b	601	.....
	Rat_2a	601	.....
	Dog_2a	601	.....
35	Cat_2a	601	.....
	Pig_2a	601	.....
	Pig_2b	601	.....
	Human_2b	601	.....
	Human_2c	601	.....
40	Human_2a	601	.....
	Rabbit_2a	601	.....
	Rabbit_2b	601	.....
45	Mouse_2a	661	SPSAQRDACLNRCAFARVEPSHKSKIVEFLQSFDEITAMTGDGVNDAPALKSEIGIAMG
	Mouse_2b	661	.....
	Rat_2b	661	.....
	Rat_2a	661	.....
	Dog_2a	661	.....
	Cat_2a	661	.....
50	Pig_2a	661	N.....E.....
	Pig_2b	661	N.....E.....
	Human_2b	661	N.....
	Human_2c	661	N.....
	Human_2a	661	N.....
55	Rabbit_2a	661	N.....
	Rabbit_2b	661	N.....

Mouse_2a	721	SGTAVAKTASEMVLADDNFSTIVAAVEEGRAIYNNMKQFIRYLISSNVGEVVCIFLTAAL
Mouse_2b	721	.....
Rat_2b	721	.....
Rat_2a	721	.....
5 Dog_2a	721	.....
Cat_2a	721	.....
Pig_2a	721	.....
Pig_2b	721	.....
Human_2b	721	.....
10 Human_2c	721	.....
Human_2a	721	.....
Rabbit_2a	721	.....
Rabbit_2b	721	.....
15 Mouse_2a	781	GFPEALIPVQLLWVNLVTDGLPATALGFNPPDLDIMNKPPRNPKEPLISGWLFFRYLAIG
Mouse_2b	781	.....
Rat_2b	781	.....
Rat_2a	781	.....
Dog_2a	781	.....
20 Cat_2a	781	.....
Pig_2a	781	.....
Pig_2b	781	.....
Human_2b	781	.....
Human_2c	781	.....
25 Human_2a	781	.....
Rabbit_2a	781	.....
Rabbit_2b	781	.....
Mouse_2a	841	CYVGAATVGAAWWFIAADGGPRVSFYQLSHFLQCKEDNPFDGVDCAIFESPYPMTMAL
30 Mouse_2b	841	.....
Rat_2b	841	..... E
Rat_2a	841	..... E
Dog_2a	841	..... D E
Cat_2a	841	..... D E
35 Pig_2a	841	..... T E V
Pig_2b	841	..... T E V
Human_2b	841	..... E
Human_2c	841	..... E
Human_2a	841	..... E
40 Rabbit_2a	841	..... E
Rabbit_2b	841	..... E
Mouse_2a	901	SVLVTIEMCNALNSLSENQSLLRMPPWENIWLGVGSICLMSMSLHFILYVEPLPLIFQITP
Mouse_2b	901	.....
45 Rat_2b	901	.....
Rat_2a	901	.....
Dog_2a	901	.....
Cat_2a	901	.....
Pig_2a	901	.....
50 Pig_2b	901	.....
Human_2b	901	.....
Human_2c	901	.....
Human_2a	901	.....
Rabbit_2a	901	.....
55 Rabbit_2b	901	.....

Mouse_2a	961	LNL TQWLMVLKISLPVILMDETLKFVARNYLEQPAILE-----
Mouse_2b	961	.....GKECVQPATKSSCSLSACTDGISWPF
Rat_2b	961	.....GKECA.....P.....
Rat_2a	961	.....AILE
5 Dog_2a	961	.....AILE
Cat_2a	961	.....AILE
Pig_2a	961	.....AILE
Pig_2b	961	.....GKEC.....-..F.....
Human_2b	961	..V.....GKEC.....-..F.....
10 Human_2c	961	..V.....VLSSL
Human_2a	961	..V.....AILE
Rabbit_2a	961	..V.....AILE
Rabbit_2b	961	..V.....GKEC....PQ.-...W...E.V....
15 Mouse_2b	1021	VLLIMPLVVWVYSTDTNFSDMFWS
Rat_2b	1020	.....
Pig_2b	1019	.....
Human_2b	1019	.....
Rabbit_2b	1019	....V...M.....LL..

**Fig. 3 Targeting construct for Serca2 flox gene modification.** Sequence information.

----- Serca2 gene

### LoxP site 1: Intron 1

5 underlined sequence = loxP site and cloning sequence

### Exon I (partial)

10 CACGGGGCTGAGCTGGAGCAGGTCAAGAACGCTCAAGGAGAGATGGGGCTCCAACGGtaggtgcggggcgc  
ccggggctgcagcggcgcggcgccccgagcgcacaaggaaatggctgaccggctccacctcgctggg  
gcttggctcggcgcggcgccccgacggctgcgagagggccggcggtccacgcgcgggtctggccatcgccg  
accttaggggtctcgaaatcaagcttatacgataccgtcgatcggacctcgagggggggggccggtagccggg  
gatcaattcgagctcgccccgggatcgatccggaaacccttaatATAACTCGTATAATGTATGCTATACG  
AAGTTATTtaggtccctcgacctcgagcccaagctccGGGGAtctcgagccggtgaccttccggccggcg  
ctcagcgagtccgattggggggggggagagggagtgggagggagggttctgcggctggctg  
agtccccccggatttatgaggcgatgttggatggaaaccctcgaccgttcttgtgcctccaaa  
15 gttgcacatctggcagaagtgtatgaccagctgaaatgactgcattgttccggagggccggagagggctt  
cggcagttcccgaggccactgattaccaggcgatgaaataatttctcggggtatcaaagtggagacagatt  
gttgcacatcatacacctataatccggcattcagacaacgatgggtgtgaatttagcagtttaataaa  
aqqcctaatacataatcttcattttcttc

----- Serca2 gene

## 20 LoxP site 2: Intron 3 5' of genomic XcmI site

underlined sequence = loxPsite, cloning sites and partial HSV-TK

ccaaattttattcttagaacattgtattcttatactgtgataggaagtaaaaatcatacagttacttgc  
ttaggtttcacaaaactgataactgtatggttcaattatgtattcacacggttaagtctgaccaggGG  
GATCCgaaacccttaatATAACTTCGTATAATGTATGCTATACGAAGTTTtaggtccctcgacctgcag  
cccaagctgatcccttagtgcagccccagctgggtcttccgcctcagaagccatagagcccaccgcac  
cccagcatgcctgctattgtcttcccaatcctcccccttgcgcctgcccccccccccccccccccc  
aatgacacactcagacaatgcgatgcatttcctcattttatttagaaaggacagtgggagttggcac  
cttccagggtcaaggaaggcacgggggaggggcaaacaacagatggctggcaactagaaggc  
ggctgatcagcgagctctagctagagaattgatccctcagaagaactcgtcaagaaggcgatagaaggc  
gatgcgtgcgaatcgggagcggcgat\*ccgtaaagcacgaggaagcgg\*cagcccattcgcgc  
tctttcagcaatatcacgggtagccaacgcgtatgt\*ctgataagcggccgcacacccaa\*ccggca  
caagtc\*atgaaatcca\*aaaaagcggccattttccacc\*atgatttt\*cggcaagcaaggc  
cattqqqgtcaccqac\*aqa\*catt\*tccgt\*c\*qqcattqgcgc\*ccct

----- HSV-TK Neo antibiotics cassette

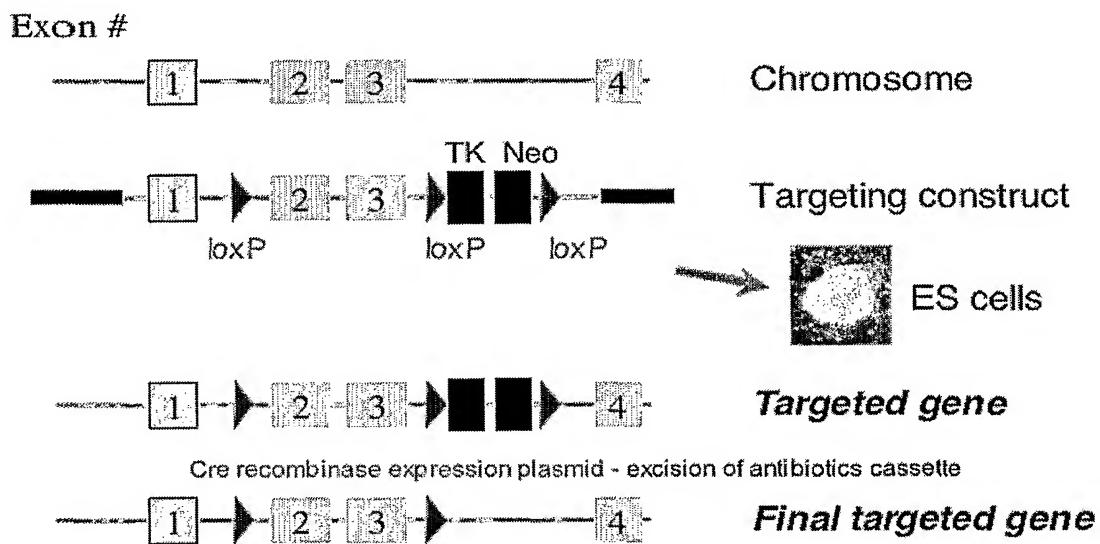
35 LoxP site 3: Intron 3 5' of genomic XcmI site

underlined sequence = loxP site, cloning sites and partial Neo gene

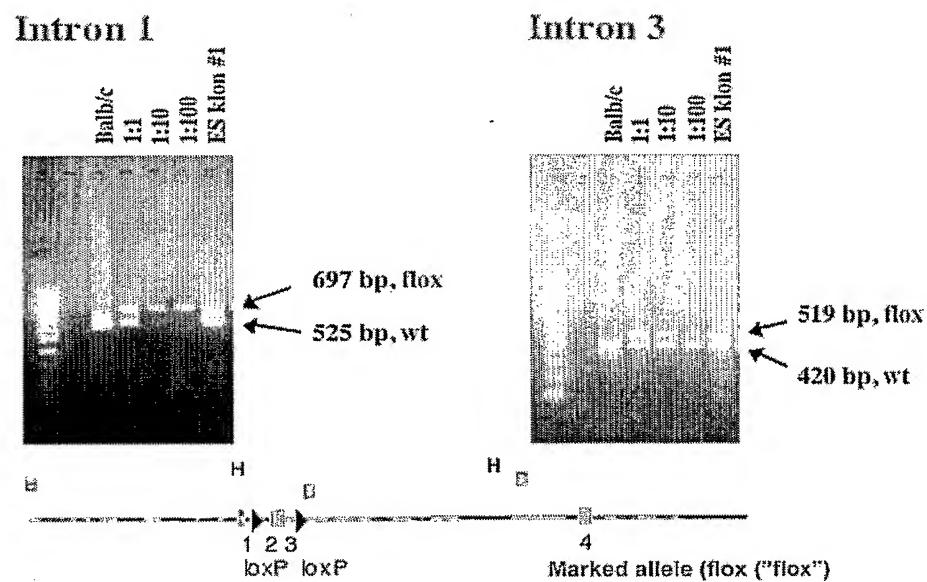
40 gttttcat\*accaccgggtcccgcc\*gatat\*ttcaccttgc\*ag\*cggtgtgtgtgtaaatg  
ttcgcgattgtttcgaaagcccc\*agcacccgcccagaatgcacatcggtcggtacgtacgcataatcgtc  
gcccgaaccaggccaccagcaagttgcgtgggtgggtttccccatcc\*gtggggac\*gtctatataaa  
acc\*gcagtagcgtggcatttctgctccggcgacttccgtggcttgcgtccggcgagggcgcaa  
cgccgtacgtcggttgcataggccgcgagaacgcgcagccgtcgaaacgcagacgcgtgtatggccgg  
ggtacgaagccatacgcgcattacaaggcgctggccgaagaggtgcggaggtttcacgcaccaagatct  
gcccgcacgcgtgttgcgcgtttaagcgggtcgctgcagggtcgctcggtgttcgaggccacacgcgtcacc  
ttaatatgcgaagtggacctcgaccgcggccggactgcacatctgcgtgttcgaattgcacaatgacaa  
45 gacgcgtggcggtttgcgcacattgggtggaaacattccaggcctgggtggagaggctttgtttcc  
tcttgc当地accacactgcgcacattgggtggaaacattccaggcctgggtggagaggctttgtttcc  
tcttgc当地aaaaccacactgcgcacattgggtggaaacattccaggcctgggtggagaggctttgtttcc  
GTTATtagtccctcgaccctgcagcccaagctgatcccgaaacccttaatATAACTTCGTATAATGTATGCTATACGAA  
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**Fig. 4A Schematic representation of genetic manipulation.**

### Serca2 (*atp2a2*) gene modification

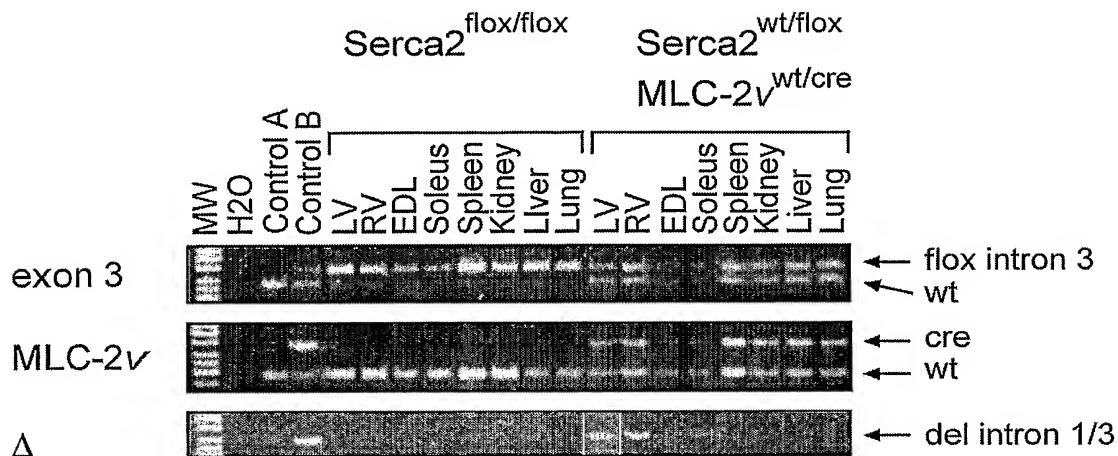


**Fig 4 B: Verification of Serca locus targeting events offspring from chimeric mice.**



**Fig. 5 Specificity of gene deletion in a test model.**

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Control A = wt ES cells (14.1a)

Control B = Upper panel: Serca2<sup>wt/flox</sup> ES cells

Middle panel: Left ventricle from MLC-2V-Cre mice

Lower panel: Serca2<sup>wt/del</sup> ES cells

LV = heart left ventricle

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RV = heart right ventricle

EDL = extensor digitorum longus muscle (fast-twitch skeletal muscle)

Soleus = soleus muscle (slow-twitch skeletal muscle)

Other tissues as indicated.

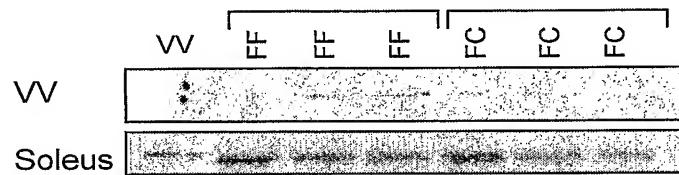
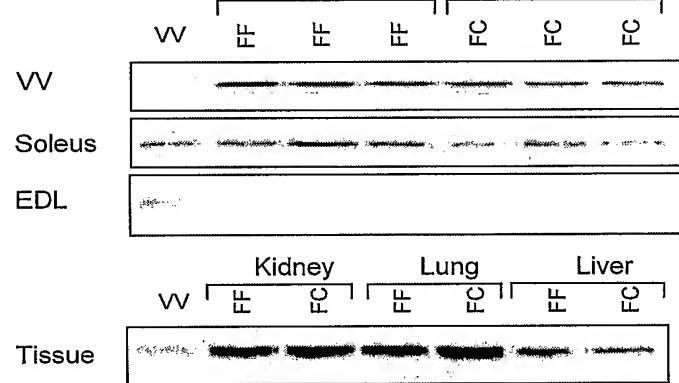
**Fig. 6 Cardiac ANP mRNA expression.**

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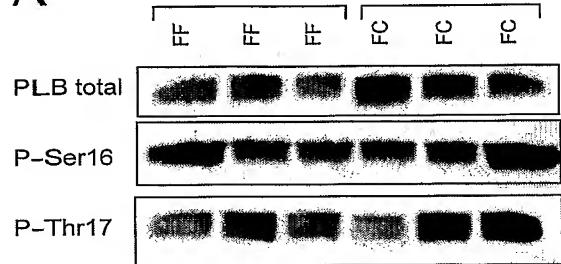


**Fig. 7 Serca2 protein expression.**

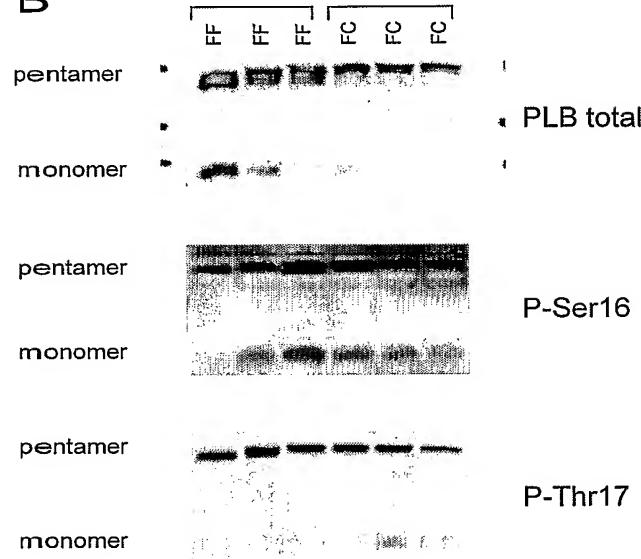
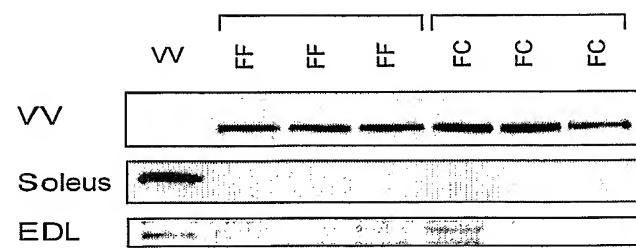
5

**A****B**

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**Fig. 8 Compensatory mechanisms in Serca<sup>flx</sup> MLC-2v-Cre mice.****A**

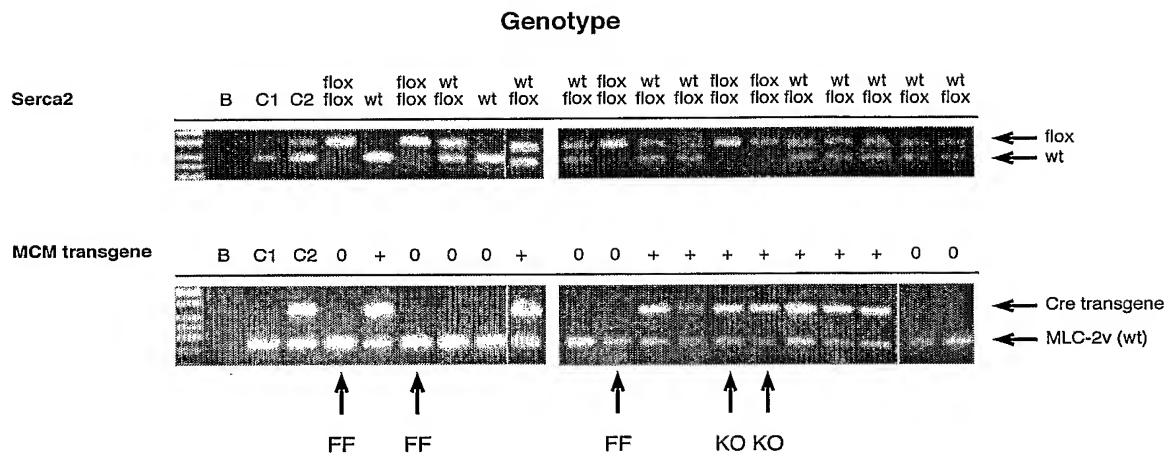
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**B****C**

10

**Figure 9      Genotypes PCR**

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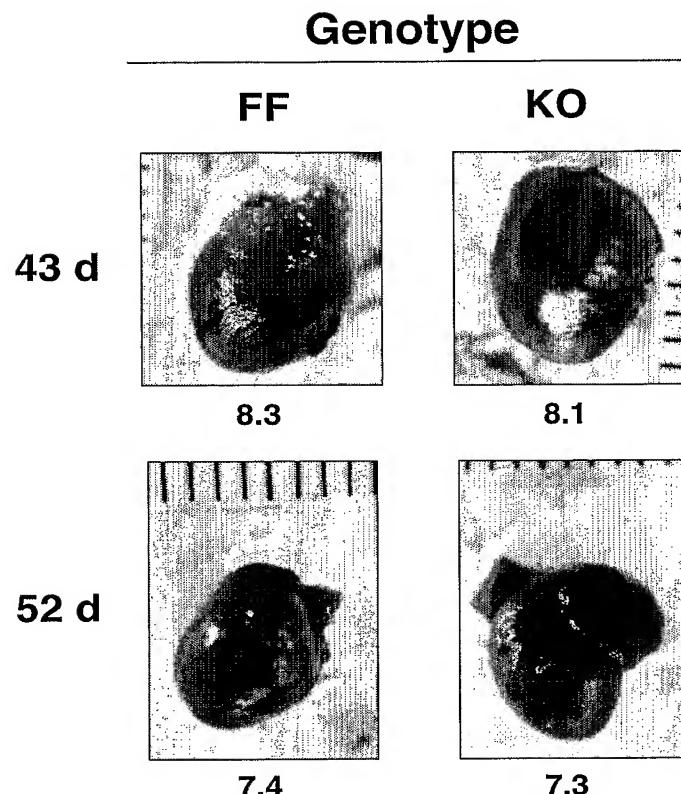


10    Generation of animals with  $\text{Serca2}^{\text{flox}}$  and MCM transgene alleles.

Genotypes FF.  $\text{Serca2}^{\text{flox/flox}}$ ;   KO,  $\text{Serca2}^{\text{flox/flox}}$  MCM

**Figure 10****Heart morphology**

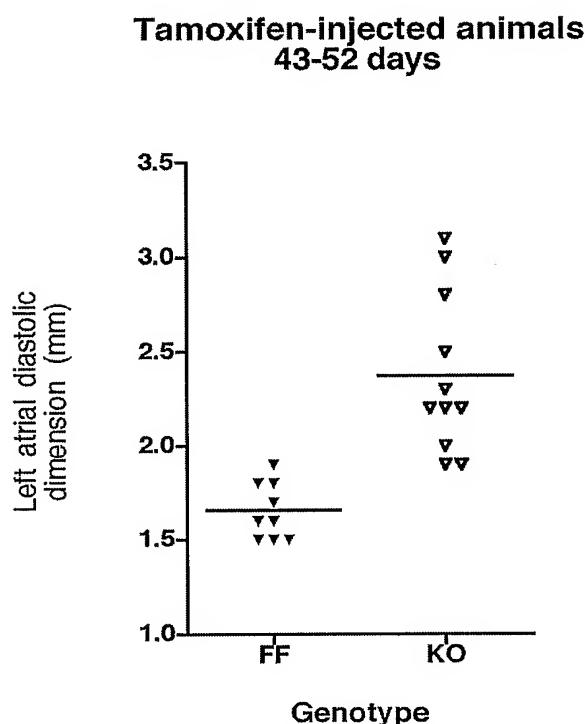
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**Figure 11 Pilot series left atrial diastolic diameter.**

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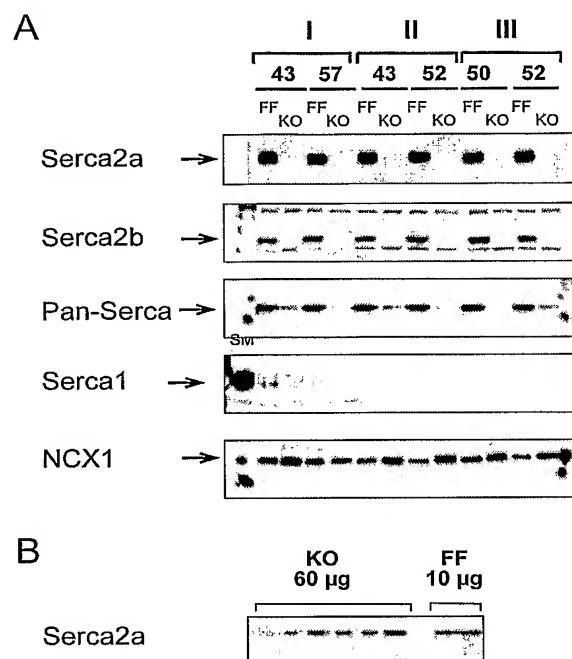


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**Figure 12** Serca protein content in tamoxifen-induced FF and KO mice

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